

APPROVED	O.G. FIG.	CLASS/SUBCLASS
	BY	
DRAFTSMAN		

FORWARD SEQUENCING PRIMER:

5'-CACTCACGCAAACGG-linker-TGTAAAACGACGGCCAGT-3'

⏟

Capture complement

⏟

Sequencing primer

REVERSE SEQUENCING PRIMER:

5'-AACTCTCCCAAGAGCAC-linker-CAGGAAACAGCTATGACC-3'

⏟

Capture complement

⏟

Sequencing primer

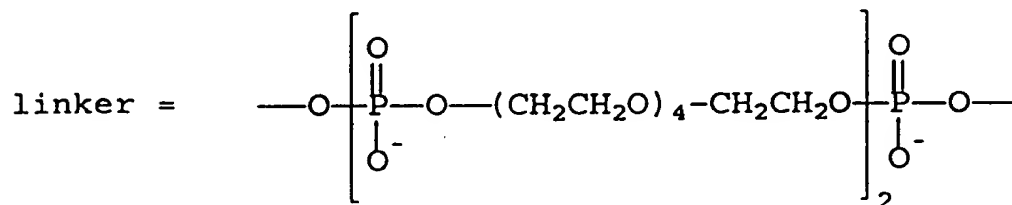
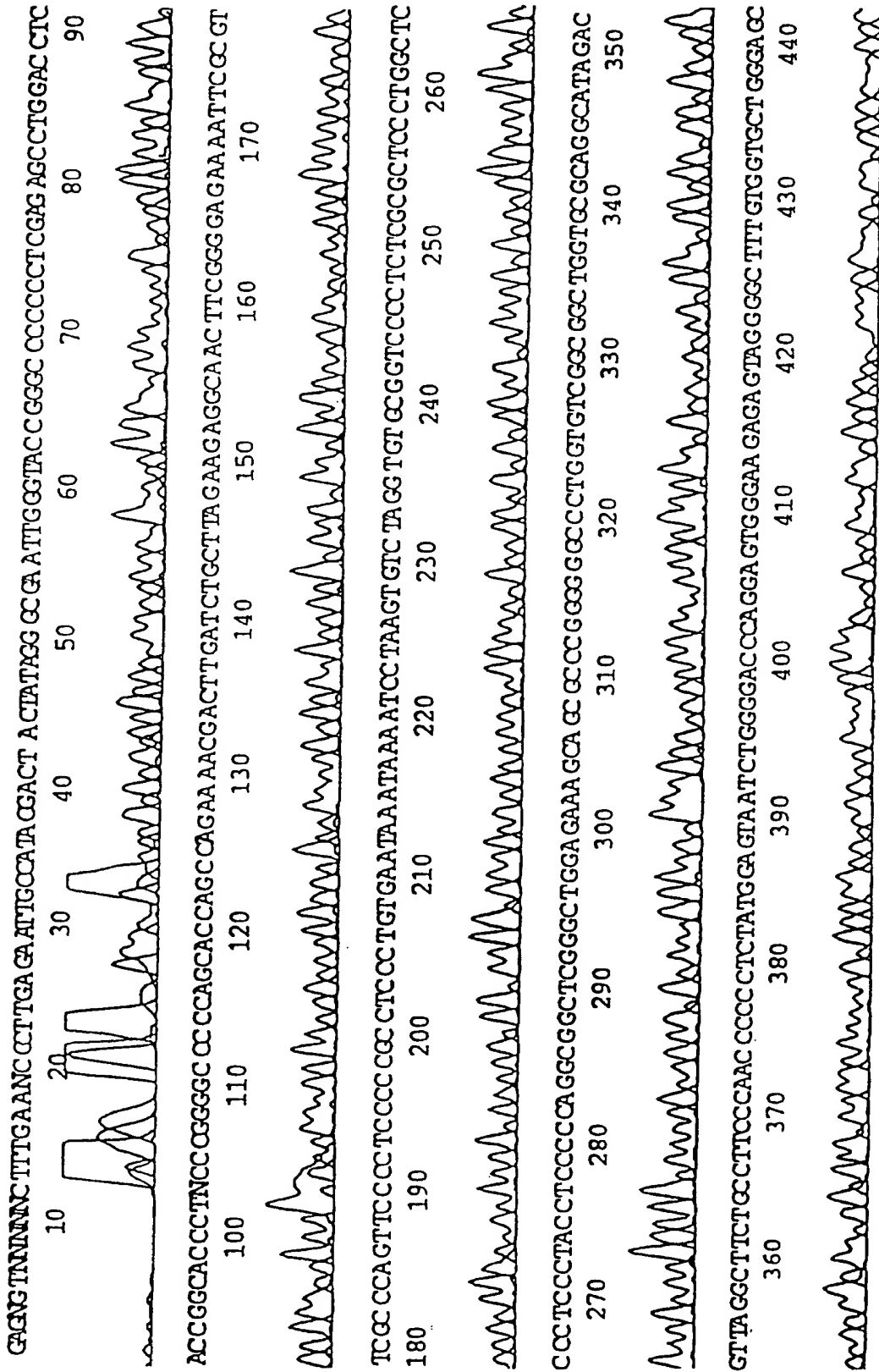


FIG. 1

APPROVED	O.G. FIG.
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FIG. 2A1

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CGAGGAAGAATGAAATGTCGCA GTTGA GTGTGTT CCTCGCATCC CAGGAGCAATGCA GTGCTGATTGTC TGTTCGTGAGATA CCCTTTTGT
450 460 470 480 490 500 510 520 530

TCGGAAACATA AGGTTTAAAGAA ACACACACAGTTT TCTCGGGTC TCTTGTTAGATGTAA ATGTCCGCCA TCCA GACCC TTACTA TCGA GTGTA
540 550 560 570 580 590 600 610 620

ACTCAATTG GGTTACTAA TCACCCCG CCCACCCGCA ACAATCAGGA TCA TTGGA ATGGGA ACCTCA TTCAA CT TCTAGGCTGAGAGTA
630 640 650 660 670 680 690 700 710

TTGCCTTCA ACCTATTTCATTCTAGACTAPG CACCAATTCAACT GAGA TAA TAAAGCAACAA TAA CCNIT GATCA TGA TGIC TA CCAA AAA
720 730 740 750 760 770 780 790 800

ATGIGCCCCGGAAAAGTCTTTTA GGAATTGTAA TGAGAAAANA AA TGAAAATGGGGCA TAAATTC TCTT
810 820 830 840 850 860 870

FIG. 2A2

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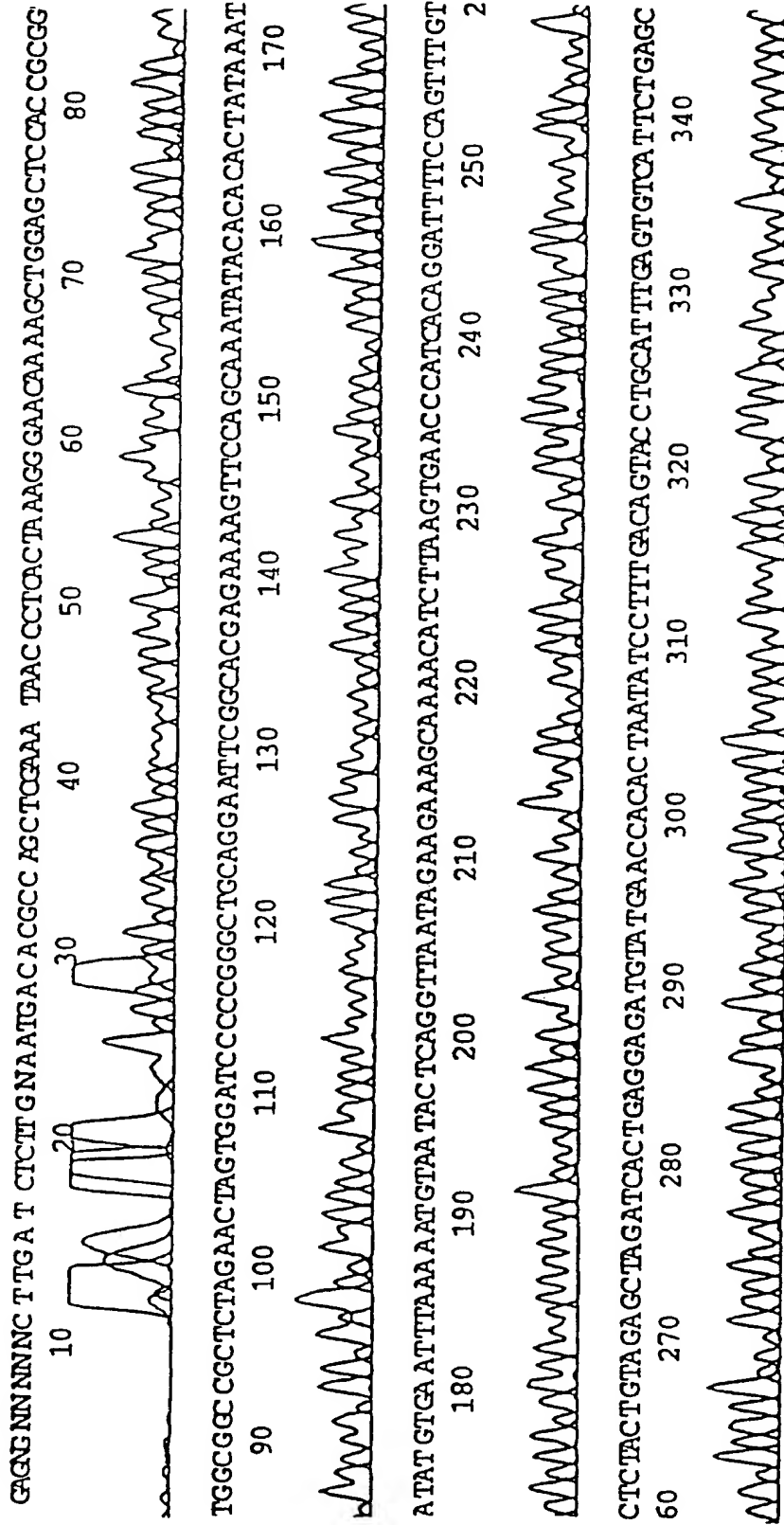


FIG. 2B1

APPROVED	O.G. FIG.	
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AAATTGAGTAA TTTTACTCAATGAGGTCA TCA TC TTCAC TAGC TTTT GGGTT TTTT TCTTTCITTTT TCTTTT TTTT TTTTAGG G
 350 360 370 380 390 400 410 420 430



GAAAGGGTGGTACA TAA GGGAC TTCTTA GTAAC TTTATT GAAGTCAA AATCATGACCCTTA TACTGCAATTTTATG GAAGCATTTAA AAAAT
 440 450 460 470 480 490 500 510 520



TATCTTGICAC TAATTGCA TTA TTGTAA GAAC TACCTCAGATTTTGTGA GTA ATATCAAAGAAAAGTC TA GTTCA TTAACAGCTACTA
 530 540 550 560 570 580 590 600 610



AGTTCTAGATNGT GTTTT GAA ACCCAGGAGTTTGGAA CAATA TATTATTA GTTTAATGGTTTA TTAATAATTTTAACTTTAA GTTCAAAC
 620 630 640 650 660 670 680 690 700 7



FIG. 2B2

APPROVED	O.G. FIG.	
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TAAAA AACAA TA TTGAAAA ANTAA ATTGAA TGGG TTA CA TTTTATGAAAAAACA TCC TG ATA GCCCTT CAAACCTTCTTT CCTTTAAAG
10 720 730 740 750 760 770 780 790



CAT TTGGGCC TACA TCTCC TTAAC TTTTCTANCOONIECCTA CCTT TGGC TACTGATTTTTTIN
800 810 820 830 840 850 860



FIG. 2B3